[Figure 1]

NCE4	MRSSPLLRSAVVAALPVLALAADGKSTRYWD	OCCKPSCGWAK	21
	*****	*****	
STCE1	MRSSPVLRTALAAALPLAALAADGKSTRYWI	CCKPSCSWPG	20
	* ****	*****	
NCE5	MQLPLTTLLTLLPALAAAQSGSGRTTRYWI	OCCKPSCAWPG	23
	Signal peptide→ Catalytic domai	<u>in</u> →	
NCE4	E4 KAPVNQPVFSCNANFQRLTDF-DAKSGCEPGGVAYSCADQTPWAVNDD		
	** ***** * **** * **** ** ***	******* *	
STCE1			
OTOLI	* ** * * * * ****		68
NCE5		A INNSVAWAS	71
NOLO	KGFAFYKTODKNON EI DUGNYKOUDKUUKTINOODK	OI WAYODDEA	,,
NCE4	4 FGFAATSIAGSNEAGWCCACYELTFTSGPVAGKKMVVQSTSTGGDLGSNH		
NOL4	****** * *** *** ********** ********		120
STCE1			
STOLI	** ** * * ** *** *** ********* * ** ****		
NOTE			121
NCE5	YGWAAVNIAGSNERQWCCACYELTFTSGPVAGKRMIVQASNTGGDLGNNH		
NOT 4	FDLN1PGGGVG1FDGCTPQFGGLPGQRYGG1SSRNE	CUDEDUVI KDG	167
NCE4	•	** ** ****	107
0.7.05.4	4 444 44444444444444444444444444444444	•••••	165
STCE1			165
	ተለ ተተቀተቀተ ተተተተ ተ ተ ተ ተ ተ ተ ተ ተ ተ ተ ተ	** ** ****	474
NCE5	FDIAMPGGGVGIFNACTDQYGAPPNGWGQRYGGISQRHE	CDAFPEKLKPG	171

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[Figure 2]

NCE4	CYWRFDWFKNADNPSFSFRQVQCPAELVARTGCRRNDDGNFPAVQIPSSS **********************************		
STCE1	CYWRFDWFKNADNPTFTFRQVQCPSELVARTGCR	RNDDGNFPVFTPPSGG	215
NCE5	CYWRFDWFLNADNPSVNWRQVSCPAEIVAKSGCS	R Linker→	206
	TSSPVGQPTSTSTTSTSTTSSPPVQPTTPS	** * ***	255
STCE1	QSSSSSSSSAKPTSTSTSTTSTKATSTTSTASS	QTSSSTGGGCAAQRWA	265
NCE5		CBD→	
NCE4	CQCGGNGWSGCTTCVAGSTCTKINDWYHQCL ***** * ********	286	
STCE1	CQCGGIGFSGCTTCVSGTTCNKQNDWYSQCL	295	
NCE5			